RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/577,973_
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RAW SEQUENCE LISTING
PATENT APPLICATION: **US/10/577,973**

DATE: 05/15/2006
TIME: 10:02:37

Input Set : **A:\50318.013001.txt**
Output Set: **N:\CRF4\05152006\J577973.raw**

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 3 <110> APPLICANT: Ensoli, Barbara
 4        Caputo, Antonella
 5        Laus, Michele
 6        Tondelli, Luisa
 7        Sparnacci, Katia
 9 <120> TITLE OF INVENTION: Nanoparticles for Delivery of a Pharmacologically Active
Agent
11 <130> FILE REFERENCE: 50318/013001
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/577,973
C--> 13 <141> CURRENT FILING DATE: 2006-05-03
13 <150> PRIOR APPLICATION NUMBER: PCT/EP2004/012420
14 <151> PRIOR FILING DATE: 2004-11-03
16 <150> PRIOR APPLICATION NUMBER: GB 0325625.2
17 <151> PRIOR FILING DATE: 2003-11-03
19 <160> NUMBER OF SEQ ID NOS: 40
21 <170> SOFTWARE: PatentIn version 3.3
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 309
25 <212> TYPE: DNA
26 <213> ORGANISM: Human immunodeficiency virus
29 <220> FEATURE:
30 <221> NAME/KEY: CDS
31 <222> LOCATION: (1)..(309)
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36 1               5                   10                  15
38 cag cct aaa act gct tgt acc aat tgc tat tgt aaa aag tgt tgc ttt      96
39 Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
40              20                  25                  30
42 cat tgc caa gtt tgt ttc ata aca aaa gcc tta ggc atc tcc tac ggc     144
43 His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly
44          35                  40                  45
46 agg aag aag cgg aga cag cgt cga aga cct cct caa ggc agt cag act     192
47 Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
48      50                  55                  60
50 cat caa gtt tct cta tca aag caa ccc acc tcc caa tcc cga ggg gac     240
51 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp
52 65                  70                  75                  80
54 ccg aca ggc ccg aag gaa cag aag aag aag gtg gag aga gag aca gag     288
55 Pro Thr Gly Pro Lys Glu Gln Lys Lys Lys Val Glu Arg Glu Thr Glu
56              85                  90                  95
58 aca gat ccg gtc cat cag tga                                         309
59 Thr Asp Pro Val His Gln
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60              100
63 <210> SEQ ID NO: 2
64 <211> LENGTH: 102
65 <212> TYPE: PRT
66 <213> ORGANISM: Human immunodeficiency virus
68 <400> SEQUENCE: 2
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71 1               5                   10                  15
74 Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
75              20                  25                  30
78 His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly
79          35                  40                  45
82 Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
83      50                  55                  60
86 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp
87 65                  70                  75                  80
90 Pro Thr Gly Pro Lys Glu Gln Lys Lys Lys Val Glu Arg Glu Thr Glu
91                  85                  90                  95
94 Thr Asp Pro Val His Gln
95              100
98 <210> SEQ ID NO: 3
99 <211> LENGTH: 261
100 <212> TYPE: DNA
101 <213> ORGANISM: Human immunodeficiency virus
104 <220> FEATURE:
105 <221> NAME/KEY: CDS
106 <222> LOCATION: (1)..(261)
108 <400> SEQUENCE: 3
109 atg gag cca gta gat cct cgt cta gag ccc tgg aag cat cca gga agt        48
110 Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser
111 1               5                   10                  15
113 cag cct aaa act gct tgt acc aat tgc tat tgt aaa aag tgt tgc ttt        96
114 Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
115              20                  25                  30
117 cat tgc caa gtt tgt ttc ata aca aaa gcc tta ggc atc tcc tac ggc        144
118 His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly
119          35                  40                  45
121 agg aag aag cgg aga cag cgt cga aga cct cct caa ggc agt cag act        192
122 Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
123      50                  55                  60
125 cat caa gtt tct cta tca aag caa ccc acc tcc caa tcc cga ggg gac        240
126 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp
127 65                  70                  75                  80
129 ccg aca ggc ccg aag gaa tag                                            261
130 Pro Thr Gly Pro Lys Glu
131                  85
134 <210> SEQ ID NO: 4
135 <211> LENGTH: 86
136 <212> TYPE: PRT
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146             20                  25                  30
149 His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly
150         35                  40                  45
153 Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
154     50                  55                  60
157 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp
158 65                  70                  75                  80
161 Pro Thr Gly Pro Lys Glu
162                 85
165 <210> SEQ ID NO: 5
166 <211> LENGTH: 261
167 <212> TYPE: DNA
168 <213> ORGANISM: Human immunodeficiency virus
171 <220> FEATURE:
172 <221> NAME/KEY: CDS
173 <222> LOCATION: (1)..(261)
175 <400> SEQUENCE: 5
176 atg gag cca gta gat cct aga cta gag ccc tgg aag cat cca gga agt        48
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178 1               5                   10                  15
180 cag cct aaa act gct ggt acc aat tgc tat tgt aaa aag tgt tgc ttt        96
181 Gln Pro Lys Thr Ala Gly Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
182             20                  25                  30
184 cat tgc caa gtt tgt ttc ata aca aaa gcc tta ggc atc tcc tat ggc       144
185 His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly
186         35                  40                  45
188 agg aag aag cgg aga cag cga cga aga cct cct caa ggc agt cag act       192
189 Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
190     50                  55                  60
192 cat caa gtt tct cta tca aag cag ccc acc tcc caa tcc cga ggg gac       240
193 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp
194 65                  70                  75                  80
196 ccg aca ggc ccg aag gaa tag                                           261
197 Pro Thr Gly Pro Lys Glu
198                 85
201 <210> SEQ ID NO: 6
202 <211> LENGTH: 86
203 <212> TYPE: PRT
204 <213> ORGANISM: Human immunodeficiency virus
206 <400> SEQUENCE: 6
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209 1               5                   10                  15
212 Gln Pro Lys Thr Ala Gly Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
213             20                  25                  30
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Input Set : **A:\50318.013001.txt**
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216 His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly
217         35              40              45
220 Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
221         50              55              60
224 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp
225 65              70              75              80
228 Pro Thr Gly Pro Lys Glu
229                 85
232 <210> SEQ ID NO: 7
233 <211> LENGTH: 261
234 <212> TYPE: DNA
235 <213> ORGANISM: Human immunodeficiency virus
238 <220> FEATURE:
239 <221> NAME/KEY: CDS
240 <222> LOCATION: (1)..(261)
242 <400> SEQUENCE: 7
243 atg gag cca gta gat cct aga cta gag ccc tgg aag cat cca gga agt      48
244 Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser
245 1               5               10              15
247 cag cct aaa act gct tgt acc aat tgc tat tgt aaa aag tgt tgc ttt      96
248 Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
249         20              25              30
251 cat tgc caa gtt tgt ttc ata aca gct gcc tta ggc atc tcc tat ggc     144
252 His Cys Gln Val Cys Phe Ile Thr Ala Ala Leu Gly Ile Ser Tyr Gly
253         35              40              45
255 agg aag aag cgg aga cag cga cga aga cct cct caa ggc agt cag act     192
256 Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
257         50              55              60
259 cat caa gtt tct cta tca aag cag ccc acc tcc caa tcc cga ggg gac     240
260 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp
261 65              70              75              80
263 ccg aca ggc ccg aag gaa tag                                         261
264 Pro Thr Gly Pro Lys Glu
265                 85
268 <210> SEQ ID NO: 8
269 <211> LENGTH: 86
270 <212> TYPE: PRT
271 <213> ORGANISM: Human immunodeficiency virus
273 <400> SEQUENCE: 8
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276 1               5               10              15
279 Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
280         20              25              30
283 His Cys Gln Val Cys Phe Ile Thr Ala Ala Leu Gly Ile Ser Tyr Gly
284         35              40              45
287 Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
288         50              55              60
291 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp
292 65              70              75              80
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295 Pro Thr Gly Pro Lys Glu
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299 <210> SEQ ID NO: 9
300 <211> LENGTH: 252
301 <212> TYPE: DNA
302 <213> ORGANISM: Human immunodeficiency virus
305 <220> FEATURE:
306 <221> NAME/KEY: CDS
307 <222> LOCATION: (1)..(252)
309 <400> SEQUENCE: 9
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312 1               5                   10                  15
314 cag cct aaa act gct tgt acc aat tgc tat tgt aaa aag tgt tgc ttt      96
315 Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
316               20                  25                  30
318 cat tgc caa gtt tgt ttc ata aca aaa gcc tta ggc atc tcc tat ggc     144
319 His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly
320               35                  40                  45
322 agg aag aag cgg aga cag cga cga aga cct cct caa ggc agt cag act     192
323 Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
324       50                  55                  60
326 cat caa gtt tct cta tca aag cag ccc acc tcc caa tcc ccg aca ggc     240
327 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Pro Thr Gly
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337 <212> TYPE: PRT
338 <213> ORGANISM: Human immunodeficiency virus
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346 Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
347               20                  25                  30
350 His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly
351               35                  40                  45
354 Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
355       50                  55                  60
358 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Pro Thr Gly
359 65                  70                  75                  80
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367 <211> LENGTH: 252
368 <212> TYPE: DNA
369 <213> ORGANISM: Human immunodeficiency virus
372 <220> FEATURE:
373 <221> NAME/KEY: CDS
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VERIFICATION SUMMARY   DATE: 05/15/2006
PATENT APPLICATION: **US/10/577,973**   TIME: 10:02:38

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L:13 M:270 C: Current Application Number differs, Replaced Current Application No
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date